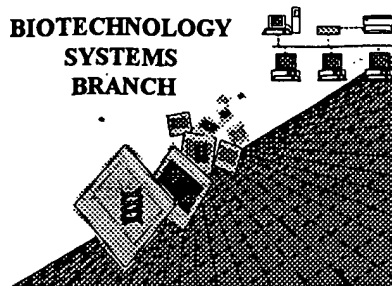


DW

RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
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The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/980,585
Source: P5/09
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

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1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission

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3. Hand Carry directly to:

U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
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4. Federal Express Delivery, 2011 South Clark Street, Crystal Plaza 2, Room 1B03-Mailroom, Box Sequence,
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PCT09

RAW SEQUENCE LISTING

DATE: 12/21/2001

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Input Set : A:\2328-124.ST25.txt

Output Set: N:\CRF3\12212001\I980585.raw

P.6
Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: LILIUS, Esa-Matti
4 VIRTÄ, Marko
6 <120> TITLE OF INVENTION: A Method to Enable Assessment of Growth and Death of Micro-organisms

8 <130> FILE REFERENCE: 2328-124

OK > 10 <140> CURRENT APPLICATION NUMBER: US/09/980,585
OK > 10 <141> CURRENT FILING DATE: 2001-12-04

10 <150> PRIOR APPLICATION NUMBER: PCT/FI00/00507

11 <151> PRIOR FILING DATE: 2000-06-07

13 <150> PRIOR APPLICATION NUMBER: FI 991296

14 <151> PRIOR FILING DATE: 1999-06-07

16 <160> NUMBER OF SEQ ID NOS: 4

18 <170> SOFTWARE: PatentIn version 3.0

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21 <211> LENGTH: 5051

22 <212> TYPE: DNA

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26 <223> OTHER INFORMATION: pGFP+Luc* plasmid

28 <220> FEATURE:

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30 <222> LOCATION: (95)..(199)

31 <223> OTHER INFORMATION: lac promoter

34 <220> FEATURE:

35 <221> NAME/KEY: CDS

36 <222> LOCATION: (289)..(1005)

37 <223> OTHER INFORMATION: GFP

40 <220> FEATURE:

41 <221> NAME/KEY: CDS

42 <222> LOCATION: (1044)..(2693)

43 <223> OTHER INFORMATION: firefly luciferase

46 <220> FEATURE:

47 <221> NAME/KEY: CDS

48 <222> LOCATION: (3251)..(4108)

49 <223> OTHER INFORMATION: beta-lactamase

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57 tcaactcatta ggcaccccag gctttacact ttatgcttcc ggctcgtatg ttgtgtggaa 180

59 ttgtgagcgg ataacaattt cacacaggaa acagctatga ccatgattac gccaaagcttg 240

61 catgcctgca ggtcgactct agaggatccc cgggtaccgg tcgccacc atg gtg agc 297

62 Met Val Ser

63 1

65 aag ggc gag gag ctg ttc acc ggg gtg gtg ccc atc ctg gtc gag ctg 345

66 Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val Glu Leu

67 5 10 15

69 gac ggc gac gta aac ggc cac aag ttc agc gtg tcc ggc gag ggc gag 393

70 Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu Gly Glu

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| 71 | 20 | 25 | 30 | 35 | |
| 73 | ggc gat gcc acc tac ggc aag ctg acc ctg aag ttc atc tgc acc acc | 441 | | | |
| 74 | Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys Thr Thr | | | | |
| 75 | 40 45 50 | | | | |
| 77 | ggc aag ctg ccc gtg ccc tgg ccc acc ctc gtg acc acc ctg acc tac | 489 | | | |
| 78 | Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Leu Thr Tyr | | | | |
| 79 | 55 60 65 | | | | |
| 81 | ggc gtg cag tgc ttc agc cgc tac ccc gac cac atg aag cag cac gac | 537 | | | |
| 82 | Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln His Asp | | | | |
| 83 | 70 75 80 | | | | |
| 85 | ttc ttc aag tcc gcc atg ccc gaa ggc tac gtc cag gag cgc acc atc | 585 | | | |
| 86 | Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg Thr Ile | | | | |
| 87 | 85 90 95 | | | | |
| 89 | ttc ttc aag gac gac ggc aac tac aag acc cgc gcc gag gtg aag ttc | 633 | | | |
| 90 | Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe | | | | |
| 91 | 100 105 110 115 | | | | |
| 93 | gag ggc gac acc ctg gtg aac cgc atc gag ctg aag ggc atc gac ttc | 681 | | | |
| 94 | Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe | | | | |
| 95 | 120 125 130 | | | | |
| 97 | aag gag gac ggc aac atc ctg ggg cac aag ctg gag tac aac tac aac | 729 | | | |
| 98 | Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn | | | | |
| 99 | 135 140 145 | | | | |
| 101 | agc cac aac gtc tat atc atg gcc gac aag cag aag aac ggc atc aag | 777 | | | |
| 102 | Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys | | | | |
| 103 | 150 155 160 | | | | |
| 105 | gtg aac ttc aag atc cgc cac aac atc gag gac ggc agc gtg cag ctc | 825 | | | |
| 106 | Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val Gln Leu | | | | |
| 107 | 165 170 175 | | | | |
| 109 | gcc gac cac tac cag cag aac acc ccc atc ggc gac ggc ccc gtg ctg | 873 | | | |
| 110 | Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu | | | | |
| 111 | 180 185 190 195 | | | | |
| 113 | ctg ccc gac aac cac tac ctg agc acc cag tcc gcc ctg agc aaa gac | 921 | | | |
| 114 | Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser Lys Asp | | | | |
| 115 | 200 205 210 | | | | |
| 117 | ccc aac gag aag cgc gat cac atg gtc ctg ctg gag ttc gtg acc gcc | 969 | | | |
| 118 | Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val Thr Ala | | | | |
| 119 | 215 220 225 | | | | |
| 121 | gcc ggg atc act ctc ggc atg gac gag ctg tac aag taaagcggcc | 1015 | | | |
| 122 | Ala Gly Ile Thr Leu Gly Met Asp Glu Leu Tyr Lys | | | | |
| 123 | 230 235 | | | | |
| 125 | gctctagaac tagtgatcc cccgtacc atg gaa gac gcc aaa aac ata aag | 1067 | | | |
| 126 | Met Glu Asp Ala Lys Asn Ile Lys | | | | |
| 127 | 240 245 | | | | |
| 129 | aaa ggc ccg gcg cca ttc tat ccg cta gag gat gga acc gct gga gag | 1115 | | | |
| 130 | Lys Gly Pro Ala Pro Phe Tyr Pro Leu Glu Asp Gly Thr Ala Gly Glu | | | | |
| 131 | 250 255 260 | | | | |
| 133 | caa ctg cat aag gct atg aag aga tac gcc ctg gtt cct gga aca att | 1163 | | | |
| 134 | Gln Leu His Lys Ala Met Lys Arg Tyr Ala Leu Val Pro Gly Thr Ile | | | | |
| 135 | 265 270 275 | | | | |

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DATE: 12/21/2001

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| 137 | gct | ttt | aca | gat | gca | cat | atc | gag | gtg | aac | atc | acg | tac | gcg | gaa | tac | 1211 |
| 138 | Ala | Phe | Thr | Asp | Ala | His | Ile | Glu | Val | Asn | Ile | Thr | Tyr | Ala | Glu | Tyr | |
| 139 | 280 | | | | | 285 | | | | 290 | | | | | 295 | | |
| 141 | ttc | gaa | atg | tcc | gtt | cgg | ttg | gca | gaa | gct | atg | aaa | cga | tat | ggg | ctg | 1259 |
| 142 | Phe | Glu | Met | Ser | Val | Arg | Leu | Ala | Glu | Ala | Met | Lys | Arg | Tyr | Gly | Leu | |
| 143 | | | | | 300 | | | | | 305 | | | | | 310 | | |
| 145 | aat | aca | aat | cac | aga | atc | gtc | gta | tgc | agt | gaa | aac | tct | ctt | caa | ttc | 1307 |
| 146 | Asn | Thr | Asn | His | Arg | Ile | Val | Val | Cys | Ser | Glu | Asn | Ser | Leu | Gln | Phe | |
| 147 | | | | | 315 | | | | | 320 | | | | | 325 | | |
| 149 | ttt | atg | ccg | gtg | ttg | ggc | gcg | tta | ttt | atc | gga | gtt | gca | gtt | gcg | ccc | 1355 |
| 150 | Phe | Met | Pro | Val | Leu | Gly | Ala | Leu | Phe | Ile | Gly | Val | Ala | Val | Ala | Pro | |
| 151 | | | 330 | | | | | 335 | | | | | 340 | | | | |
| 153 | gcg | aac | gac | att | tat | aat | gaa | cgt | gaa | ttg | ctc | aac | agt | atg | aac | att | 1403 |
| 154 | Ala | Asn | Asp | Ile | Tyr | Asn | Glu | Arg | Glu | Leu | Leu | Asn | Ser | Met | Asn | Ile | |
| 155 | | 345 | | | | | 350 | | | | | 355 | | | | | |
| 157 | tcg | cag | cct | acc | gta | gtg | ttt | gtt | tcc | aaa | aag | ggg | ttg | caa | aaa | att | 1451 |
| 158 | Ser | Gln | Pro | Thr | Val | Val | Phe | Val | Ser | Lys | Lys | Gly | Leu | Gln | Lys | Ile | |
| 159 | 360 | | | | | 365 | | | | 370 | | | | | 375 | | |
| 161 | ttg | aac | gtg | caa | aaa | aaa | tta | cca | ata | atc | cag | aaa | att | att | atc | atg | 1499 |
| 162 | Leu | Asn | Val | Gln | Lys | Lys | Leu | Pro | Ile | Ile | Gln | Lys | Ile | Ile | Ile | Met | |
| 163 | | | | | 380 | | | | | 385 | | | | | 390 | | |
| 165 | gat | tct | aaa | acg | gat | tac | cag | gga | ttt | cag | tcg | atg | tac | acg | ttc | gtc | 1547 |
| 166 | Asp | Ser | Lys | Thr | Asp | Tyr | Gln | Gly | Phe | Gln | Ser | Met | Tyr | Thr | Phe | Val | |
| 167 | | | | 395 | | | | 400 | | | | | 405 | | | | |
| 169 | aca | tct | cat | cta | cct | ccc | ggt | ttt | aat | gaa | tac | gat | ttt | gta | cca | gag | 1595 |
| 170 | Thr | Ser | His | Leu | Pro | Pro | Gly | Phe | Asn | Glu | Tyr | Asp | Phe | Val | Pro | Glu | |
| 171 | | | 410 | | | | | 415 | | | | | 420 | | | | |
| 173 | tcc | ttt | gat | cgt | gac | aaa | aca | att | gca | ctg | ata | atg | aac | tcc | tct | gga | 1643 |
| 174 | Ser | Phe | Asp | Arg | Asp | Lys | Thr | Ile | Ala | Leu | Ile | Met | Asn | Ser | Ser | Gly | |
| 175 | | 425 | | | | | 430 | | | | | 435 | | | | | |
| 177 | tct | act | ggg | tta | cct | aag | ggt | gtg | gcc | ctt | ccg | cat | aga | act | gcc | tgc | 1691 |
| 178 | Ser | Thr | Gly | Leu | Pro | Lys | Gly | Val | Ala | Leu | Pro | His | Arg | Thr | Ala | Cys | |
| 179 | 440 | | | | | 445 | | | | 450 | | | | | 455 | | |
| 181 | gtc | aga | ttc | tcg | cat | gcc | aga | gat | cct | att | ttt | ggc | aat | caa | atc | att | 1739 |
| 182 | Val | Arg | Phe | Ser | His | Ala | Arg | Asp | Pro | Ile | Phe | Gly | Asn | Gln | Ile | Ile | |
| 183 | | | | | 460 | | | | | 465 | | | | | 470 | | |
| 185 | ccg | gat | act | gcg | att | tta | agt | gtt | gtt | cca | ttc | cat | cac | ggt | ttt | gga | 1787 |
| 186 | Pro | Asp | Thr | Ala | Ile | Leu | Ser | Val | Val | Pro | Phe | His | His | Gly | Phe | Gly | |
| 187 | | | | 475 | | | | | 480 | | | | | 485 | | | |
| 189 | atg | ttt | act | aca | ctc | gga | tat | ttg | ata | tgt | gga | ttt | cga | gtc | gtc | tta | 1835 |
| 190 | Met | Phe | Thr | Thr | Leu | Gly | Tyr | Leu | Ile | Cys | Gly | Phe | Arg | Val | Val | Leu | |
| 191 | | | 490 | | | | | 495 | | | | | 500 | | | | |
| 193 | atg | tat | aga | ttt | gaa | gaa | gag | ctg | ttt | tta | cga | tcc | ctt | cag | gat | tac | 1883 |
| 194 | Met | Tyr | Arg | Phe | Glu | Glu | Glu | Leu | Phe | Leu | Arg | Ser | Leu | Gln | Asp | Tyr | |
| 195 | | 505 | | | | | 510 | | | | | 515 | | | | | |
| 197 | aaa | att | caa | agt | gcg | ttg | cta | gta | cca | acc | cta | ttt | tca | ttc | ttc | gcc | 1931 |
| 198 | Lys | Ile | Gln | Ser | Ala | Leu | Leu | Val | Pro | Thr | Leu | Phe | Ser | Phe | Phe | Ala | |
| 199 | 520 | | | | | 525 | | | | 530 | | | | | 535 | | |
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DATE: 12/21/2001

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| 202 | Lys | Ser | Thr | Leu | Ile | Asp | Lys | Tyr | Asp | Leu | Ser | Asn | Leu | His | Glu | Ile | |
| 203 | | | | | 540 | | | | | 545 | | | | | 550 | | |
| 205 | gct | tct | ggg | ggc | gca | cct | ctt | tcg | aaa | gaa | gtc | ggg | gaa | gcg | gtt | gca | 2027 |
| 206 | Ala | Ser | Gly | Gly | Ala | Pro | Leu | Ser | Lys | Glu | Val | Gly | Glu | Ala | Val | Ala | |
| 207 | | | | 555 | | | | | 560 | | | | | 565 | | | |
| 209 | aaa | cgc | ttc | cat | ctt | cca | ggg | ata | cga | caa | gga | tat | ggg | ctc | act | gag | 2075 |
| 210 | Lys | Arg | Phe | His | Leu | Pro | Gly | Ile | Arg | Gln | Gly | Tyr | Gly | Leu | Thr | Glu | |
| 211 | | | 570 | | | | | 575 | | | | 580 | | | | | |
| 213 | act | aca | tca | gct | att | ctg | att | aca | ccc | gag | ggg | gat | gat | aaa | ccg | ggc | 2123 |
| 214 | Thr | Thr | Ser | Ala | Ile | Leu | Ile | Thr | Pro | Glu | Gly | Asp | Asp | Lys | Pro | Gly | |
| 215 | | 585 | | | | | 590 | | | | 595 | | | | | | |
| 217 | gcg | gtc | ggt | aaa | gtt | gtt | cca | ttt | ttt | gaa | gcg | aag | gtt | gtg | gat | ctg | 2171 |
| 218 | Ala | Val | Gly | Lys | Val | Val | Pro | Phe | Phe | Glu | Ala | Lys | Val | Val | Asp | Leu | |
| 219 | 600 | | | | | 605 | | | | 610 | | | | | 615 | | |
| 221 | gat | acc | ggg | aaa | acg | ctg | ggc | gtt | aat | cag | aga | ggc | gaa | tta | tgt | gtc | 2219 |
| 222 | Asp | Thr | Gly | Lys | Thr | Leu | Gly | Val | Asn | Gln | Arg | Gly | Glu | Leu | Cys | Val | |
| 223 | | | | 620 | | | | | 625 | | | | | 630 | | | |
| 225 | aga | gga | cct | atg | att | atg | tcc | ggt | tat | gta | aac | aat | ccg | gaa | gcg | acc | 2267 |
| 226 | Arg | Gly | Pro | Met | Ile | Met | Ser | Gly | Tyr | Val | Asn | Asn | Pro | Glu | Ala | Thr | |
| 227 | | | 635 | | | | | 640 | | | | | 645 | | | | |
| 229 | aac | gcc | ttg | att | gac | aag | gat | gga | tgg | cta | cat | tct | gga | gac | ata | gct | 2315 |
| 230 | Asn | Ala | Leu | Ile | Asp | Lys | Asp | Gly | Trp | Leu | His | Ser | Gly | Asp | Ile | Ala | |
| 231 | | | 650 | | | | 655 | | | | | 660 | | | | | |
| 233 | tac | tgg | gac | gaa | gac | gaa | cac | ttc | ttc | ata | gtt | gac | cgc | ttg | aag | tct | 2363 |
| 234 | Tyr | Trp | Asp | Glu | Asp | Glu | His | Phe | Phe | Ile | Val | Asp | Arg | Leu | Lys | Ser | |
| 235 | | 665 | | | | 670 | | | | 675 | | | | | | | |
| 237 | tta | att | aaa | tac | aaa | gga | tac | cag | gtg | gcc | ccc | gct | gaa | ttg | gag | tcg | 2411 |
| 238 | Leu | Ile | Lys | Tyr | Lys | Gly | Tyr | Gln | Val | Ala | Pro | Ala | Glu | Leu | Glu | Ser | |
| 239 | 680 | | | | 685 | | | | 690 | | | | | | 695 | | |
| 241 | ata | ttg | tta | caa | cac | ccc | aac | atc | ttc | gac | gcg | ggc | gtg | gca | ggt | ctt | 2459 |
| 242 | Ile | Leu | Leu | Gln | His | Pro | Asn | Ile | Phe | Asp | Ala | Gly | Val | Ala | Gly | Leu | |
| 243 | | | 700 | | | | | 705 | | | | | | 710 | | | |
| 245 | ccc | gac | gat | gac | gcc | ggt | gaa | ctt | ccc | gcc | gcc | gtt | gtt | gtt | ttg | gag | 2507 |
| 246 | Pro | Asp | Asp | Asp | Ala | Gly | Glu | Leu | Pro | Ala | Ala | Val | Val | Val | Leu | Glu | |
| 247 | | | 715 | | | | | 720 | | | | | 725 | | | | |
| 249 | cac | gga | aag | acg | atg | acg | gaa | aaa | gag | atc | gtg | gat | tac | gtc | gcc | agt | 2555 |
| 250 | His | Gly | Lys | Thr | Met | Thr | Glu | Lys | Glu | Ile | Val | Asp | Tyr | Val | Ala | Ser | |
| 251 | | | 730 | | | | 735 | | | | | 740 | | | | | |
| 253 | caa | gta | aca | acc | gcc | aaa | aag | ttg | cgc | gga | gga | gtt | gtg | ttt | gtg | gac | 2603 |
| 254 | Gln | Val | Thr | Thr | Ala | Lys | Lys | Leu | Arg | Gly | Gly | Val | Val | Phe | Val | Asp | |
| 255 | | 745 | | | | 750 | | | | | 755 | | | | | | |
| 257 | gaa | gta | ccg | aaa | ggt | ctt | acc | gga | aaa | ctc | gac | gca | aga | aaa | atc | aga | 2651 |
| 258 | Glu | Val | Pro | Lys | Gly | Leu | Thr | Gly | Lys | Leu | Asp | Ala | Arg | Lys | Ile | Arg | |
| 259 | 760 | | | | 765 | | | | 770 | | | | | 775 | | | |
| 261 | gag | atc | ctc | ata | aag | gcc | aag | aag | ggc | gga | aag | tcc | aaa | ttg | | | 2693 |
| 262 | Glu | Ile | Leu | Ile | Lys | Ala | Lys | Lys | Gly | Gly | Lys | Ser | Lys | Leu | | | |
| 263 | | | 780 | | | | | 785 | | | | | | | | | |
| 265 | taaaatgtaa | ctgtatttcag | cgatgacgaa | attcttagct | attgtaatac | tctaggggct | | | | | | | | | | | 2753 |
| 267 | gcaggaattc | gatatcaagc | ttatcgatac | cgtcgacctc | gagggggggc | ccttttcgtct | | | | | | | | | | | 2813 |

RAW SEQUENCE LISTING

DATE: 12/21/2001

PATENT APPLICATION: US/09/980,585

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Output Set: N:\CRF3\12212001\I980585.raw

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277 ttaagggcct cgtgatacgc ctatttttat aggttaatgt catgataata atggtttctt 3113
279 agacgtcagg tggcactttt cggggaaatg tgcgcggaac ccctatttgt ttatttttct 3173
281 aaatacatc aaatatgtat ccgctcatga gacaataacc ctgataaatg cttcaataat 3233
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284 Met Ser Ile Gln His Phe Arg Val Ala Leu Ile
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287 ccc ttt ttt gcg gca ttt tgc ctt cct gtt ttt gct cac cca gaa acg 3331
288 Pro Phe Phe Ala Ala Phe Cys Leu Pro Val Phe Ala His Pro Glu Thr
289 805 810 815
291 ctg gtg aaa gta aaa gat gct gaa gat cag ttg ggt gca cga gtg ggt 3379
292 Leu Val Lys Val Lys Asp Ala Glu Asp Gln Leu Gly Ala Arg Val Gly
293 820 825 830
295 tac atc gaa ctg gat ctc aac agc ggt aag atc ctt gag agt ttt cgc 3427
296 Tyr Ile Glu Leu Asp Leu Asn Ser Gly Lys Ile Leu Glu Ser Phe Arg
297 835 840 845
299 ccc gaa gaa cgt ttt cca atg atg agc act ttt aaa gtt ctg cta tgt 3475
300 Pro Glu Glu Arg Phe Pro Met Ser Thr Phe Lys Val Leu Leu Cys
301 850 855 860
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304 Gly Ala Val Leu Ser Arg Ile Asp Ala Gly Gln Glu Gln Leu Gly Arg
305 865 870 875 880
307 cgc ata cac tat tct cag aat gac ttg gtt gag tac tca cca gtc aca 3571
308 Arg Ile His Tyr Ser Gln Asn Asp Leu Val Glu Tyr Ser Pro Val Thr
309 885 890 895
311 gaa aag cat ctt acg gat ggc atg aca gta aga gaa tta tgc agt gct 3619
312 Glu Lys His Leu Thr Asp Gly Met Thr Val Arg Glu Leu Cys Ser Ala
313 900 905 910
315 gcc ata acc atg agt gat aac act gcg gcc aac tta ctt ctg aca acg 3667
316 Ala Ile Thr Met Ser Asp Asn Thr Ala Ala Asn Leu Leu Leu Thr Thr
317 915 920 925
319 atc gga gga ccg aag gag cta acc gct ttt ttg cac aac atg ggg gat 3715
320 Ile Gly Gly Pro Lys Glu Leu Thr Ala Phe Leu His Asn Met Gly Asp
321 930 935 940
323 cat gta act cgc ctt gat cgt tgg gaa ccg gag ctg aat gaa gcc ata 3763
324 His Val Thr Arg Leu Asp Arg Trp Glu Pro Glu Leu Asn Glu Ala Ile
325 945 950 955 960
327 cca aac gac gag cgt gac acc acg atg cct gta gca atg gca aca acg 3811
328 Pro Asn Asp Glu Arg Asp Thr Thr Met Pro Val Ala Met Ala Thr Thr
329 965 970 975
331 ttg cgc aaa cta tta act ggc gaa cta ctt act cta gct tcc cgg caa 3859
332 Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu Thr Leu Ala Ser Arg Gln
333 980 985 990
335 caa tta ata gac tgg atg gag gcg gat aaa gtt gca gga cca ctt ctg 3907
336 Gln Leu Ile Asp Trp Met Glu Ala Asp Lys Val Ala Gly Pro Leu Leu
337 995 1000 1005

```

09/980,585 6

<210> 2
<211> 239
<212> PRT
<213> Artificial

see item 11 on Error Summary Sheet

<400> 2

↓
The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/980,585

DATE: 12/21/2001

TIME: 13:17:19

Input Set : A:\2328-124.ST25.txt

Output Set: N:\CRF3\12212001\I980585.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:23 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:395 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:397 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:397 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:462 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:464 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:464 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:609 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:611 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:611 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: